

## RAW SEQUENCE LISTING ERROR REPORT

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Application Serial Number:	09/991,518
Source:	0185
Date Processed by STIC:	3-7-03

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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

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Revised 01/29/2002



## Does Not Comply Corrected Diskette Needed

OIPE

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/991,518

DATE: 03/07/2003

TIME: 13:26:00

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Output Set: N:\CRF4\03072003\I991518.raw

1 <110> APPLICANT: Fenical, William Jensen,

W--> 2 <120> TITLE OF INVENTION: Marine Actinomycete Taxon for Drug and

W--> 3 <130> FILE REFERENCE: 6627-PA1022<150> 60/249,356<151> 2000-11-

C--> 4 <140> CURRENT APPLICATION NUMBER: US/09/991,518

C--> 4 <141> CURRENT FILING DATE: 2003-02-19

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## ERRORED SEQUENCES

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22

Please Contact Bob wax (703) 308-4216

Patentla formatting error

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/09/991,518

DATE: 03/07/2003 TIME: 13:26:01

Input Set : A:\pto.vsk.txt

Output Set: N:\CRF4\03072003\I991518.raw

## Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:0; Line(s) 1,3

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/991,518

DATE: 03/07/2003 TIME: 13:26:01

Input Set : A:\pto.vsk.txt

Output Set: N:\CRF4\03072003\I991518.raw

L:2 M:283 W: Missing Blank Line separator, <120> field identifier L:3 M:283 W: Missing Blank Line separator, <130> field identifier L:4 M:270 C: Current Application Number differs, Replaced Current Application No L:4 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:4 M:283 W: Missing Blank Line separator, <160> field identifier L:5 M:282 E: Numeric Field Identifier Missing, <210> is required. L:5 M:282 E: Numeric Field Identifier Missing, <211> is required. L:5 M:282 E: Numeric Field Identifier Missing, <212> is required. L:5 M:282 E: Numeric Field Identifier Missing, <213> is required. L:5 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQ ID NO L:5 M:283 W: Missing Blank Line separator, <400> field identifier L:10 M:282 E: Numeric Field Identifier Missing, <211> is required. L:10 M:282 E: Numeric Field Identifier Missing, <212> is required. L:10 M:282 E: Numeric Field Identifier Missing, <213> is required. L:10 M:200 E: Mandatory Header Field missing, <400> is required. L:4 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (2) Counted (1)